RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	10/534,74213
Source:	TFWP
Date Processed by STIC:	07/31/2006

ENTERED



IFWP

RAW SEQUENCE LISTING DATE: 07/31/2006 PATENT APPLICATION: US/10/534,742B TIME: 14:14:03

Input Set : A:\10534742.txt

Output Set: N:\CRF4\07312006\J534742B.raw

```
3 <110> APPLICANT: Corrado FOGHER
     5 <120> TITLE OF INVENTION: Food flours with specific technological
characteristics and low
             allergenicity
     8 <130> FILE REFERENCE: 4161-12 / BX1898R
    10 <140> CURRENT APPLICATION NUMBER: US 10/534,742B
                                                             (Pg-6)
    11 <141> CURRENT FILING DATE: 2005-05-12
    13 <150> PRIOR APPLICATION NUMBER: PCT/IB2003/005092
    14 <151> PRIOR FILING DATE: 2003-11-12
    16 <150> PRIOR APPLICATION NUMBER: IT BO2002A000714
    17 <151> PRIOR FILING DATE: 2002-11-13
    19 <160> NUMBER OF SEO ID NOS: 44
    21 <170> SOFTWARE: MS Word
    23 <210> SEQ ID NO: 1
    24 <211> LENGTH: 830
    25 <212> TYPE: PRT
    26 <213> ORGANISM: Wheat
    28 <400> SEQUENCE: 1
    30 Met Thr Lys Arg Leu Val Leu Phe Ala Ala Val Val Ala Leu Val
                                           10
    33 Ala Leu Thr Ala Ala Glu Gly Glu Ala Ser Gly Gln Leu Gln Cys Glu
                   20
                                       25
    36 Arg Glu Leu Gln Glu His Ser Leu Lys Ala Cys Arg Gln Val Val Asp
                                   40
              35
    39 Gln Gln Leu Arg Asp Val Ser Pro Glu Cys Gln Pro Val Gly Gly Gly
    42 Pro Val Ala Arg Gln Tyr Glu Gln Gln Val Val Pro Pro Lys Gly
                                               75
    45 Gly Ser Phe Tyr Pro Gly Glu Thr Thr Pro Pro Gln Gln Leu Gln Gln
    48 Ser Ile Leu Trp Gly Ile Pro Ala Leu Leu Arg Arg Tyr Tyr Leu Ser
                   100
                                       105
                                                           110
    51 Val Thr Ser Pro Gln Gln Val Ser Tyr Tyr Pro Gly Gln Ala Ser Ser
                                   120
    54 Gln Arg Pro Gly Gln Gly Gln Fro Gly Gln Gly Gln Glu Tyr
                               135
    57 Tyr Leu Thr Ser Pro Gln Gln Ser Gly Gln Trp Gln Gln Pro Gly Gln
                           150
                                               155
    60 Gly Gln Ala Gly Tyr Tyr Pro Thr Ser Pro Gln Gln Ser Gly Gln Glu
                       165
                                           170
    63 Gln Pro Gly Tyr Tyr Pro Thr Ser Pro Trp Gln Pro Glu Gln Leu Gln
                   180
                                       185
    66 Gln Pro Thr Gln Gly Gln Gln Arg Gln Gln Pro Gly Gln Gly Gln Gln
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200

195

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69 Leu Arg Gln Gly Gln Gln Gln Gln Ser Gly Gln Gly Gln Pro Arg 215 72 Tyr Tyr Pro Thr Ser Ser Gln Gln Pro Gly Gln Leu Gln Gln Leu Ala 230 235 75 Gln Gly Gln Gln Gln Gln Pro Glu Arg Gly Gln Gln Gln Gln Gln Gln 245 250 78 Ser Gly Gln Gly Gln Leu Gly Gln Gly Gln Gly Gln Gly Fro 265 81 Gly Gln Lys Gln Gln Ser Gly Gln Gly Gln Gly Tyr Tyr Pro Ile 280 84 Ser Pro Gln Gln Leu Gly Gln Gly Gln Gln Ser Gly Gln Gly Gln Leu 295 87 Gly Tyr Tyr Pro Thr Ser Pro Gln Gln Ser Gly Gln Gly Gln Ser Gly 310 315 90 Tyr Tyr Pro Thr Ser Ala Gln Gln Pro Gly Gln Leu Gln Gln Ser Thr 325 330 93 Gln Glu Gln Leu Gly Gln Glu Gln Asp Gln Gln Ser Gly Gln 340 345 96 Gly Arg Gln Gly Gln Ser Gly Gln Arg Gln Gln Asp Gln Gln Ser 355 360 365 99 Gly Gln Gly Gln Pro Gly Gln Arg Gln Pro Gly Tyr Tyr Ser Thr 375 102 Ser Pro Gln Gln Leu Gly Gln Gly Gln Pro Arg Tyr Tyr Pro Thr Ser 390 395 105 Pro Gln Gln Pro Gly Gln Glu Gln Pro Arg Gln Leu Gln Gln Pro 108 Glu Gln Gly Gln Gly Gln Gln Pro Glu Gln Gly Gln Gly Gln 420 425 111 Gln Pro Gly Gln Gly Gln Gly Gln Gln Pro Gly Gln Gln Gln 440 114 Gly Gln Gln Pro Gly Gln Gly Gln Pro Gly Tyr Tyr Pro Thr Ser Pro 455 460 117 Gln Gln Ser Gly Gln Gly Gln Pro Gly Tyr Tyr Pro Thr Ser Pro Gln 470 475 120 Gln Ser Gly Gln Leu Gln Gln Pro Ala Gln Gly Gln Gln Pro Gly Gln 485 490 123 Glu Gln Gln Gly Gln Pro Gly Gln Gly Gln Gln Gly Gln Pro 500 505 126 Gly Gln Gly Gln Pro Gly Gln Gly Gln Pro Gly Tyr Tyr Pro Thr 520 129 Ser Pro Gln Gln Ser Gly Gln Glu Gln Leu Glu Gln Trp Gln Gln 535 540 132 Ser Gly Gln Gly Gln Pro Gly His Tyr Pro Thr Ser Pro Leu Gln Pro 555 135 Gly Gln Gly Gln Pro Gly Tyr Tyr Pro Thr Ser Pro Gln Gln Ile Gly 570 565 138 Gln Gly Gln Gln Pro Gly Gln Leu Gln Gln Pro Thr Gln Gly Gln Gln 585 141 Gly Gln Gln Pro Gly Gln Gly Gln Gly Gln Gln Pro Gly Gln Gly

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PATENT APPLICATION: US/10/534,742B TIME: 14:14:03

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142			595					600					605			
	Gln	Gln		Gln	Gln	Pro	Glv		Glv	Gln	Gln	Pro		Gln	Glv	Gln
145		610	- 1				615		011	U	01	620	017	0	U -1	0211
			Tvr	Tvr	Pro	Thr		Len	Gln	Gln	Ser		Gln	Glv	Gln	Gln
	625	1	-1-	-7-		630				Ÿ 	635		02	0-1	0	640
		Glv	Gln	Tro	Gln		Pro	Glv	Gln	Glv		Pro	Glv	Tur	Tur	
151					645	01	110	Q_Y	0111	650	шец		Ory	- 7 -	655	110
	Thr	Ser	Ser	Len	Gln	Pro	G] 11	Gln	Glv		Gln	Glv	Tvr	Tur		Thr
154				660			01 u	Q	665	0.111	01	O _T	- 1 -	670	110	
	Ser	Gln	Gln		Pro	Glv	Gln	Glv		Gln	Pro	Glv	Gln		Gln	Gln
157			675			V-1		680				017	685			02
	Ser	Glv		Glv	Gln	Gln	Glv		Tvr	Pro	Thr	Ser		Gln	Gln	Ser
160		690		1			695	-1-	-1-			700		U	V	001
	Glv		Glv	Gln	Gln	·Pro		Gln	Trp	Leu	Gln		Glv	Gln	Trp	Leu
	705		2			710	1				715		V-1	V		720
		Ser	Glv	Tvr	Tyr		Thr	Ser	Pro	Gln		Leu	Glv	Gln	Glv	
166			1	-1-	725					730			1		735	
	Gln	Pro	Arq	Gln	Trp	Leu	Gln	Pro	Ara		Glv	Gln	Gln	Glv		Tvr
169				740	Ľ				745		2			750	- 4 -	-2-
171	Pro	Thr	Ser	Pro	Gln	Gln	Ser	Glv	Gln	Glv	Gln	Gln	Leu	Glv	Gln	Glv
172			755					760		4			765	. 4		
174	Gln	Gln	Gly	Tyr	Tyr	Pro	Thr	Ser	Pro	Gln	Gln	Ser	Gly	Gln	Gly	Gln
175		770	-	•	•		775					780	•		•	
177	Gln	Gly	Tyr	Asp	Ser	Pro	Tyr	His	Val	Ser	Ala	Glu	His	Gln	Ala	Ala
178	785	_	_	_		790	_				795					800
180	Ser	Leu	Lys	Val	Ala	Lys	Ala	Gln	Gln	Leu	Ala	Ala	Gln	Leu	Pro	Ala
181					805					810					815	
183	Met	Cys	Arg	Leu	Glu	Gly	Gly	Asp	Ala	Leu	Leu	Ala	Ser	Gln		
184				820					825					830		
187	<210)> SI	EQ II	ON C	: 2											
188	<21	l> LI	ENGTI	I: 8.	15											
189	<212	2> T	YPE:	PRT												
190	<213	3 > OI	RGAN:	ISM:	Whea	at										
192	<400)> SI	EQUE	ICE:	2											
194	Met	Thr	Lys	Arg	Leu	Val	Leu	Phe	Ala	Ala	Val	Val	Val	Ala	Leu	Val
195					5					10					15	
197	Ala	Leu	Thr	Ala	Ala	Glu	Gly	Glu	Ala	Ser	Gly	Gln	Leu	Gln	Cys	Glu
198				20					25					30		
	Arg			Gln	Glu	His	Ser	Leu	Lys	Ala	Cys	Arg	Gln	Val	Val	Asp
201	_		35			_		40					45			
			Leu	Arg	Asp	Val		Pro	Glu	Cys	Gln	Pro	Val	Gly	Gly	Gly
204		50					55					60				
		Val	Ala	Arg	Gln	_	Glu	Gln	Gln	Val		Val	Pro	Pro	Lys	
207		_				70	_		_		75	_	_		_	80
	Gly	Ser	Phe	Tyr	Pro	Gly	Glu	Thr	Thr		Pro	Gln	Gln	Leu		Gln
210	_		_	_	85		_		_	90	_	_	_		95	_
	Ser	Ile	Leu		Gly	IIe	Pro	Ala		Leu	Arg	Arg	Tyr		Leu	Ser
213		-1	_	100	~7	~7		_	105	_	_			110	_	_
215	Val	Thr	Ser	Pro	Gln	Gln	Val	Ser	Tyr	Tyr	Pro	Gly	Gln	Ala	Ser	Ser

RAW SEQUENCE LISTING
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216			115					120					125			
				Glv	Gln	Glv	Gln		Glu	ጥ ን/	ጥኒንን	T.e.11		Ser	Pro	Gln
219		130	110	013	01	013	135	0111	Olu	- y -	- y -	140	1111	001	110	0111
			Glv	Gln	Trp	Gln		Pro	Glv	Gln	Glv		Sar	Glv	Tt/r	ጥኒም
	145	001	017	0111		150	0111	110		0111	155	0111	001	O _T	- 7 -	160
		Thr	Ser	Pro	Gln		Ser	Gl v	Gln	Larg		Dro	Glv	ጥላታ	ጥኒ/ዮ	
225	110	1111	Der	110	165	GIII	Der	Gry	GIII	170	GIII	FIO	Gry	ıyı	175	FIO
	Thr	Ser	Pro	Trn	Gln	Dro	G111	Gln	T.011		Gln	Dro	Thr	Gln		Gln
228	1111	Ser	110	180	GIII	FIO	Giu	GIII	185	GIII	GIII	FIU	1111	190	Gry	GIII
	Gln	Δνα	Gln.		Pro	Glv	Gln	G1 17		Gln	T.011	λνα	Gl n		Gln	Gln
231	GTir	n. a	195	GIII	110	Gry	GIII	200	GIII	GIII	пец	Arg	205	GLY	GIII	GIII
	Glv	Gln		Ser	Gly	Gln	Glv		Pro	Ara	ጥኒኒኒ	Туг		Thr	Ser	Ser
234	Q ₁ y	210	OIII	DCI	OLY	OIII	215	GIII	110	n-9	- Y -	220	110	1111	Der	DCI
	Gln		Pro	Glv	Gln	T.e.11		Gln	T.e.11	Δla	Gln		Gln	Gln	Glv	Gln
	225	0111	110	O _T y	0111	230	01	0111	Lcu	niu	235	OLY	0111	OIII	GLY	240
		Pro	Glu	Ara	Gly		Gln	Glv	Gln	Gln		Glv	Gln	Glv	Gln	-
240	0111	110	OIU	y	245	0111	OIII	CLY	QIII	250	DCL	OLY	GIII	CLY	255	0111
	Len	Glv	Gln	Glv	Gln	Gln	Glv	Gln	Gln		Glv	Gln	Lvs	Gln		Ser
243	200	O.J	01	260	0	01	017	0111	265	110	O _T	0111	цу	270	0111	UCI
	Glv	Gln	Glv		Gln	Glv	Tvr	Tvr		Tle	Ser	Pro	Gln		T.e11	Glv
246	- 1		275	01	· · · · ·	017	-1-	280			501		285	01		
	Gln	Glv		Gln	Ser	Glv	Gln		Gln	Len	Glv	Tvr		Pro	Thr	Ser
249	01	290		0111	001	017	295	O+1			017	300	-1-			501
	Pro		Gln	Ser	Gly	Gln		Gln	Ser	Glv	Tvr		Pro	Thr	Ser	Ala
	305				0-1	310	0-1			0-1	315	-1-			202	320
		Gln	Pro	Glv	Gln		Gln	Gln	Ser	Thr		Glu	Gln	Gln	Leu	
255		-		- 1	325					330					335	2
	Gln	Glu	Gln	Gln	Asp	Gln	Gln	Ser	Glv		Glv	Ara	Gln	Glv		Gln
258				340	•				345		4	_		350		
260	Ser	Glv	Gln	Arq	Gln	Gln	Asp	Gln	Gln	Ser	Glv	Gln	Glv	Gln	Gln	Pro
261		•	355				-	360			•		365			
263	Gly	Gln	Arg	Gln	Pro	Gly	Tyr	Tyr	Ser	Thr	Ser	Pro	Gln	Gln	Leu	Gly
264	_	370	_			_	·375	_				380				_
266	Gln	Gly	Gln	Pro	Arg	Tyr	Tyr	Pro	Thr	Ser	Pro	Gln	Gln	Pro	Gly	Gln
267	385	_			_	390	_				395				_	400
269	Glu	Gln	Gln	Pro	Arg	Gln	Leu	Gln	Gln	Pro	Glu	Gln	Gly	Gln	Gln	Gly
270					405					410			_		415	_
272	Gln	Gln	Pro	Glu	Gln	Gly	Gln	Gln	Gly	Gln	Gln	Gln	Arg	Gln	Gly	Glu
273				420		_			425				_	430	_	
275	Gln	Gly	Gln	Gln	Pro	Gly	Gln	Gly	Gln	Gln	Gly	Gln	Gln	Pro	Gly	Gln
276			435			_		440			-		445		_	
278	Gly	Gln	Pro	Gly	Tyr	Tyr	Pro	Thr	Ser	Pro	Gln	Gln	Ser	Gly	Gln	Gly
279		450					455					460		_		_
281	Gln	Pro	Gly	Tyr	Tyr	Pro	Thr	Ser	Pro	Gln	Gln	Ser	Gly	Gln	Leu	Gln
	465		_	•		470					475		=			480
284	Gln	Pro	Ala	Gln	Gly	Gln	Gln	Pro	Gly	Gln	Glu	Gln	Gln	Gly	Gln	Gln
285					485				_	490				=	495	
287	Pro	Gly	Gln	Gly	Gln	Gln	Pro	Gly	Gln	Gly	Gln	Pro	Gly	Tyr	Tyr	Pro
288				500					505					510		

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290 Thr Ser Pro Gln Gln Ser Gly Gln Glu Gln Leu Glu Gln Trp Gln 293 Gln Ser Gly Gln Gly Gln Pro Gly His Tyr Pro Thr Ser Pro Leu Gln 535 540 296 Pro Gly Gln Gly Gln Pro Gly Tyr Tyr Pro Thr Ser Pro Gln Gln Ile 550 555 299 Gly Gln Gly Gln Pro Gly Gln Leu Gln Pro Thr Gln Gly Gln 565 570 302 Gln Gly Gln Gln Pro Gly Gln Gly Gln Gly Gln Gln Pro Gly Glu 585 305 Gly Gln Gln Gln Gln Pro Gly Gln Gly Gln Gln Pro Gly Gln Gly 308 Gln Pro Gly Tyr Tyr Pro Thr Ser Leu Gln Gln Ser Gly Gln Gly Gln 615 311 Gln Pro Gly Gln Trp Gln Gln Pro Gly Gln Gly Gln Pro Gly Tyr Tyr 630 635 314 Pro Thr Ser Ser Leu Gln Pro Glu Gln Gly Gln Gly Tyr Tyr Pro 645 650 317 Thr Ser Gln Gln Gln Pro Gly Gln Gly Pro Gln Pro Gly Gln Trp Gln 660 665 320 Gln Ser Gly Gln Gly Gln Gly Tyr Tyr Pro Thr Ser Pro Gln Gln 675 680 323 Ser Gly Gln Gly Gln Gln Pro Gly Gln Trp Leu Gln Pro Gly Gln Trp 695 326 Leu Gln Ser Gly Tyr Tyr Leu Thr Ser Pro Gln Gln Leu Gly Gln Gly 710 715 329 Gln Gln Pro Arg Gln Trp Leu Gln Pro Arg Gln Gly Gln Gly Tyr 725 730 332 Tyr Pro Thr Ser Pro Gln Gln Ser Gly Gln Gly Gln Leu Gly Gln 740 745 335 Gly Gln Gln Gly Tyr Tyr Pro Thr Ser Pro Gln Gln Ser Gly Gln Gly 755 760 338 Gln Gln Gly Tyr Asp Ser Pro Tyr His Val Ser Ala Glu His Gln Ala 775 341 Ala Ser Leu Lys Val Ala Lys Ala Gln Gln Leu Ala Ala Gln Leu Pro 790 795 344 Ala Met Cys Arg Leu Glu Gly Gly Asp Ala Leu Leu Ala Ser Gln 347 <210> SEQ ID NO: 3 348 <211> LENGTH: 839 349 <212> TYPE: PRT 350 <213> ORGANISM: Wheat 352 <400> SEQUENCE: 3 354 Met Ala Lys Arg Leu Val Leu Phe Val Ala Val Val Ala Leu Val 357 Ala Leu Thr Val Ala Glu Gly Glu Ala Ser Glu Gln Leu Gln Cys Glu 358 20 25 360 Arg Glu Leu Gln Glu Leu Gln Glu Arg Glu Leu Lys Ala Cys Gln Gln 40

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 07/31/2006
PATENT APPLICATION: US/10/534,742B TIME: 14:14:04

Input Set : A:\10534742.txt

Output Set: N:\CRF4\07312006\J534742B.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220>

to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:44; Xaa Pos. 2

VERIFICATION SUMMARY

DATE: 07/31/2006

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Input Set : A:\10534742.txt

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L:2115 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:44 after pos.:0